9889 6889 6989	0 K 126 E 179 2 128 T 180	0 F 216 F 218 F 270	K 305 N 359 S 307 K 359	0 L 395 L 397 L 449
90 KGSYGVDEMDVKSLR KGSHGVDEMKVDELL QGAPGIDGVSTDQLR KGAPGADGMTVADLA	166 RRGHNAVRQAKQWMK RSAKQAIKKAKEYME RNAHDAVRQAQGYIQ RSTHQAIEMARAHVT	256 LDELDNELEKRGLKF LTELDRELEKRGKKF LDDLDKELEKRGLKF LDELDRELERRGHRF	346 KQRIRELTSRRHSMK KAKIKALTARSNALN KQRIRQLTNPNWSIS RDRLRRLLRSVRARK	436 GYWRVAHSPIMDYAL SYWRIANSPILSTTL GAWRTIKTPQLHQAL GPWWNSGAQHMNYAL
YGVDEI SHGVDGI APGIDGY APGADGI	5 SHNAVR( AKQAIKI AHDAVR( IHQAIEI	SLONELI SLONELI SLORELI SLORELI	SIRELT: KIKAIT: KIRQLT! KIRMLL!	NEVAHSI NELANSI NETTKTI
			ı	
61 SRENLIQALERVEKN DRONMNLAYKRVKSN ARDNLITALKRVEAN APANLRRAYQRVVSN	151 YDPTESERSEGERPH FEKTESENSYGERPK FDPDFSPSSEGERPG FDPLFSDYSYGFRPG	241 255 TPQGPLSPLLSNIN CPQGPLSPLLSNIM TPQGGPLSPLLANIL TPQGGPLSPLLSNIL TVGGPLSPLLSNIL TVGGPLSPLLSNIL TVGGPLSPLLSNIL TV	331 -DPKMRVSKESVKRL GKIGIRVHEKSVKKF -KARIRLAPRSIQRL -QPRLRVARMSLDRL	421 GIKKAKAWQWANSRK GINNSKAWEFANTRK GLKETAVWEIANTRK
60 MLERIL DDGKLYETNILERIL MALLERIL TWTNAEPDTLMERVL	136 DRFLQQAIAQILTQL DRTIQQAISQVLTPI DRLIQQAILQELTPI DRLIQQALQQQLTPI	226 TGVMERGLVSPNTEG SCWINGVVSETERG AGVMIEGVKVQTEFG AGIMSGGVVSPRQEG	316 BWNRKELGESFTRGK PWRRKELGESFTRGK PWRRAELGESFTPER AWKCDYLGYGMSWHQ VII	406 EWKTTSARQKNLVRL QWKKVKTKYDNLRKF QWKRVRTRIRELRA QWKQPPTRLRNLMRL
31 	121 135 KPNGGVRKLGIPTWM KPDGGIRLLGIPTWV KPGGGTRQLGIPTWV KPQGGTRQLGIPSWV II	211 225 226 240 IQDPRVLQLIRRYLQ TGVMERGLVSPNTEG IKDRRVLKLIRLYLQ SGVMINGVVSETERG VKDRRVLKLIRAYLQ AGVMIEGVKVQTEFG IKDKCVLRLIRRYLE AGIMSGGVVSPRQEG	301 RLKLKVNREKSAVDR KLKLKVNKEKSAVDR TLKLKVNEEKSAVDR RLKLTVNRKKSQVAR VI	391 405 406 IDAWIRRILAMIRWK EWK. LDEWTRRILAMCYWK QWK IEGWIRRILALCQWL QWK LDGWVRHKLRCVIWR QWK
30  YKEGWSCEIRVELON 	106 EGSYFPKPVRRVEIP NGKYCPKAVRRVEIP AGTYRPAPVRRVGIP AGEYHPQAV <u>RAVEIP</u>	210 DKVNHDRLMRKLSSR DTVNHDKLMALVARK DRVNHDILMSRVARK DRVNHDILMSRVARK	300 AGLRIMESVTSFIEN AGDRVMRSITRFIEN AGQRVKQSIQRFLEK AGERVLVSVERFIRE	YY 376 390 LGYYQVVDTPSILAQ LNYFGIAEMTKLAKK IGYFRLVETPSVLQT ASYFKLSOSKRPLEE
1 	105 LHLHENWTSIRNEII QYLKQNGKTLIASIF DXIRAHWSTIRAQLL GYVKQYWPTLKARLL	181 EGYRWVVDIDLEKFF EGYKWVVDIDLAKYF EGYRYVVDMDLEKFF AGHRWCVELDLEKFF	271 285 286 300 VRYADDCNIYVRSKR AGLRIMESVTSFIEN CRYADDNNYYRSKK AGDRVMRSITRFIEN CRYADDANIYVRSLR AGGRVKQSIQRFLEK VRYADDANIYVRSPR AGERVLVSVERFIRE V	LN G 361 375 MSDRLRRLNRYLTGW IENRIIKLRQCIIGW MPERIHRVNQYWGW MATVIERINPVLRGW
1 1. B.halodurans 2 C.acetobutylicum -MKN 3 Bst803-2065 4 PalcaligenesMatur MPPV	91 105 1 B.halodurans LHLHENWTSIRNEII 2 C.acetobutylicum QYLKQNGKTLIASIF 3 <b>Bst803-2065</b> DYIRAHWSTIRAQLL 4 PalcaligenesMatur GYVKQYWPTLKARLL	1 B.halodurans 2 C.acetobutylicum 3 Bst803-2065 4 PalcaligenesMatur	1 B.halodurans 2 C.acetobutylicum 3 Bst803-2065 4 PalcaligenesMatur	361 1 B.halodurans MSDF 2 C.acetobutylicum IENF 3 Bst803-2065 MPEF 4 PalcaligenesMatur MPEF

Fig. 1

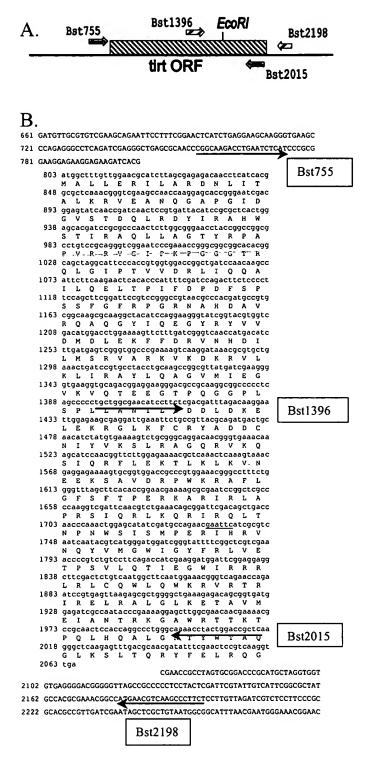


Fig. 2

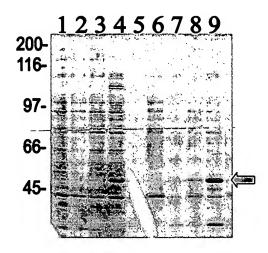


Fig. 3

## S1234567891011121314

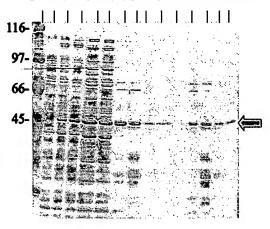


Fig. 4

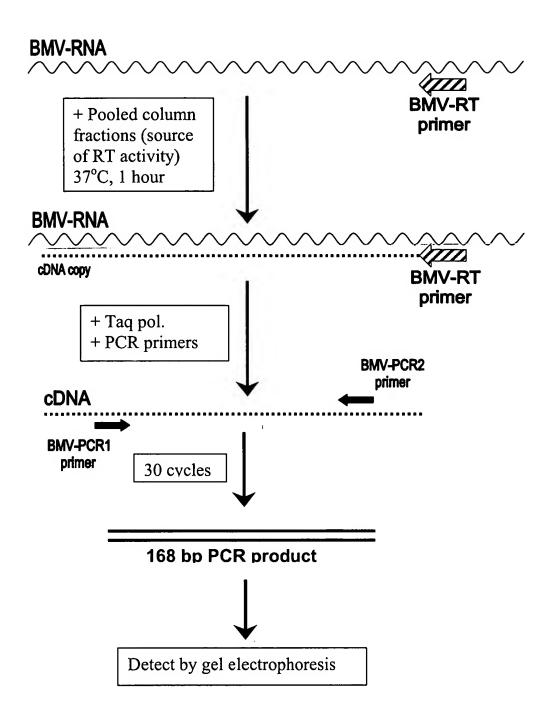


Fig. 5

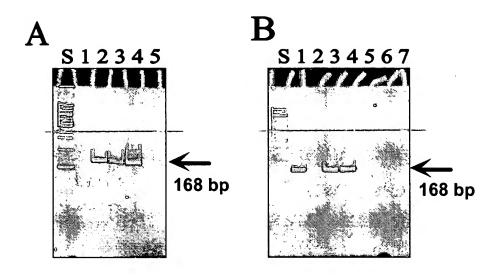


Fig. 6

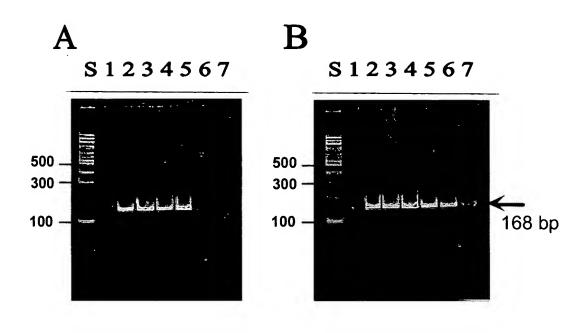
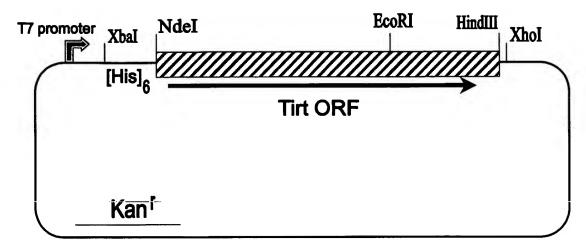


Fig. 7





Plasmid pTirt#16

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Fig. 8